DRAFTSMAH

CLASS SUBCLASS

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Figure 1A

- 2252644	C A COCCO		CACCTGAGGC	NCTCTCCACC	ACAGCCCACC
>3353644	CATTIGAGGC	CAIAIAAAGI		CCTCTCCACC	
>2804743			Gridge	CCTCTCCMCC	C
>1891065	CAMMMCACCC	C λ M λ M M λ λ C M	CACCTGAGGC	CCTCTCCACC	_
Consensus	CATTIGAGGC	CATATAAAGT	CACCIGAGGC	CCICICCACC	ACAGCCCACC
>3353644	AGTGACCATG	AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>2804743	AGTGACCATG		TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1891065	AGTGACCATG		TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1543671		AAGGCTGTGC	TGCTTGCCCT	GTTGATGGCA	GGCTTGGCCC
>1543671IH		AAGGCTGTGC		GTTGATGGCA	
Consensus		AAGGCTGTGC		GTTGATGGCA	
Consensus	Adroneeme	12.00010100			
>3353644	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>2804743	TGCAGCCAGG	CACTGCCCTG		CCTGCAAAGC	
>1891065	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>1543671	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
>1543671IH	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
Consensus	TGCAGCCAGG	CACTGCCCTG	CTGTGCTACT	CCTGCAAAGC	CCAGGTGAGC
00110011001					
>3353644	AACGAGGACT	GCCTGCAGGT		ACCCAGCTGG	
>2804743	AACGAGGACT	GCCTGCAGGT		ACCCAGCTGG	
>1891065	AACGAGGACT	GCCTGCAGGT		ACCCAGCTGG	
>1543671	AACGAGGACT	GCCTGCAGGT	GGAGAACTGC	ACCCAGCTGG	GGGAGCAGTG
>1543671IH	AACGAGGACT	GCCTGCAGGT	GGAGAACTGC	ACCCAGCTGG	GGGAGCAGTG
>1863905	CGAGGACT	GCCTGCAGGT	GGAGAACTGC	ACCCAGCTGG	GGGAGCAGTG
Consensus	AACGAGGACT	GCCTGCAGGT	GGAGAACTGC	ACCCAGCTGG	GGGAGCAGTG
>3353644	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT		ATCAGCAAAG
>2804743	CTGGACCGCG	CGCATCCGCG			ATCAGCAAAG
>1891065	CTGGACCGCG	CGCATCCGCG		CCTGACCGTC	
>1543671	CTGGACCGCG	CGCATCCGCG			ATCAGCAAAG
>1543671IH		CGCATCCGCG			ATCAGCAAAG
>1863905	CTGGACCGCG	CGCATCCGCG	CAGTTGGCCT		ATCAGCAAAG
Consensus	CTGGACCGCG	CGCATCCGCG	CAGTTGGCĊT	CCTGACCGTC	ATCAGCAAAG
			GATGANTCAC	AGGANTACTA	C
>3353644		GAACTGCGTG			CGTGGGCAAG
>2804743	GCTGCAGCTT		=		CGTGGGCAAG
>1891065	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACIACIA	CGIGGGCAAG
>1543671	GCTGCAGC			3 CO 3 CM 3 CM 3	COMCCCCAAC
>1543671IH	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACTACTA	CGTGGGCAAG
>1863905	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACTACTA	CGTGGGCAAG
Consensus	GCTGCAGCTT	GAACTGCGTG	GATGACTCAC	AGGACTACTA	CGTGGGCAAG
- 2004743	AAGAACATCA	ССТССТСТ			
>2804743		CGTGCTGTGA	CACCGACT		
>1891065	AAGAACAICA	CGTGCTGTG	CACCGACTTC	TGCAACGCCA	GCGGGGCCCA
>1543671IH	AAGAACAICA	COTOCTORO	CACCGACTTC	TGCAACGCCA	GCGGGGCCCA
>1863905	AAGAACATCA	CMCCMCMC	CACCGACTIC	TGCAACGCCA	GCGGGGCCCA
>1314679					GCGGGGCCCA
Consensus	AAGAACATCA	A CGTGCTGTGA	1 CACCGACIIC	, recuredect	. Jegoodeech

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Figure 1B

		`			
>1543671IH	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
>1863905	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
>1314679				GCTGCTCCCT	
Consensus	TGCCCTGCAG	CCGGCTGCCG	CCATCCTTGC	GCTGCTCCCT	GCACTCGGCC
>1543671IH	TGCTGCTCTG	GGGACCCGGC	CAGCTATAGG	CTCTGGGGGG	CCCCGCTGCA
>1863905	TGCTGCTCTG	GGGACCC			
>1314679	TGCTGCTCTG	GGGACCCGGC	CAGCTATAGG	CTCTGGGGGG	CCCCGCTGCA
Consensus	TGCTGCTCTG	GGGACCCGGC	CAGCTATAGG	CTCTGGGGGG	CCCCGCTGCA
>1543671IH		GGTGTGGTGC		TGTGCCACTC	
>1314679	GCCCACACTG	GGTGTGGTGC	CCCAGGCCTC	TGTGCCACTC	CTCACAGACC
>1901337					С
Consensus	GCCCACACTG	GGTGTGGTGC	CCCAGGCCTC	TGTGCCACTC	CTCACAGACC
				•	
>1543671IH				AGGCACATCC	
>1314679				AGGCACATCC	
>1901337	TGGCCCAGTG			AGGCACATCC	
>1900086				AGGCACATCC	
Consensus	TGGCCCAGTG	GGAGCCTGTC	CTGGTTCCTG	AGGCACATCC	TAACGCAAGT
>1543671IH				CACCCTGACC	
>1901337				CACCCTGACC	
>1900086				CACCCTGACC	
Consensus	CTGACCATGT	ATGTCTGCAC	CCCTGTCCCC	CACCCTGACC	CTCCCATGGC
			000101m010	CMCMA CMCA C	2 C 2 C 2 TC C C C
>1543671IH				CTCTAGTGAC	
>1901337				CTCTAGTGAC CTCTAGTGAC	
>1900086				CTCTAGTGAC	
Consensus	CCTCTCCAGG	ACTCCCACCC	GGCAGATCAG	CICIAGIGAC	ACAGATCCGC
- 15 42 671 TU	CTCCACATCC	ССССТССААС	CCTCTCTCTCT	GCTGTTTCCA	TGGCCCAGCA
>1543671IH >1901337		CCCCTCCAAC			TGGCCCAGCA
>1901337		CCCCTCCAAC			TGGCCCAGCA
Consensus		CCCCTCCAAC			TGGCCCAGCA
Consensus	CIGCAGNICO	ccccrccinic	00101010	33233333	
>1543671IH	TTCTCCACCC	TTAACCCTGT	GCTCAGGCAC	CTCTTCCCCC	AGGAAGCCTT
>1901337	TTCTCCACCC		GCTCAGGCAC		
>1901037	TTCTCCACCC				AGGAAGCCTT
>2325070					AGGAAGCCTT
Consensus	TTCTCCACCC				AGGAAGCCTT
Compensus					
>1543671IH	CCCTGCCCAC	CCCATCTATG	ACTTGAGCCA	GGTCTGGTCC	GTGGTGTCCC
>1900086	CCCTGCCCAC				
>2325070			ACTTGAGCCA	GGTCTGGTCC	GTGGTGTCCC
>3969672					GTGGTGTCCC
Consensus	CCCTGCCCAC				GTGGTGTCCC
>1543671IH					AAGGCTGAGA
>2325070	CCGCACCCAG	CAGGGGACAG	GCACTCAGGA	GGGCCCAGTA	AAGGCTGAGA
>3969672					AAGGCTGAGA
Consensus	CCGCACCCAG	CAGGGGACAG	GCACTCAGGA	GGGCCCAGTA	AAGGCTGAGA

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Figure 1C

>1543671IH	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>2325070	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>3969672	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
Consensus	TGAAGTGGAC	TGAGTAGAAC	TGGAGGACAA	GAGTCGACGT	GAGTTCCTGG
>1543671IH	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
>2325070	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
>3969672	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
Consensus	GAGTCTCCAG	AGATGGGGCC	TGGAGGCCTG	GAGGAAGGGG	CCAGGCCTCA
				•	
>1543671IH	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
>2325070	CATTCGTGGG	GCTCCCTGAA			
>3969672	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
Consensus	CATTCGTGGG	GCTCCCTGAA	TGGCAGCCTG	AGCACAGCGT	AGGCCCTTAA
>1543671IH	TAAACACCTG	TTGGATAAGC	CCA		
>3969672	TAAACACCTG	TTGGAT			
Consensus	TAAACACCTG	TTGGATAAGC	CCA		

1900086 2325070 3969672

1863905 1314679 1901337

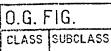
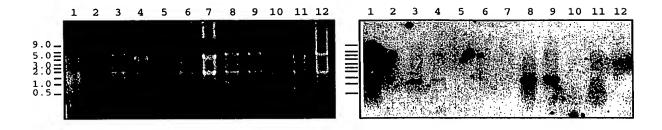




Figure 3



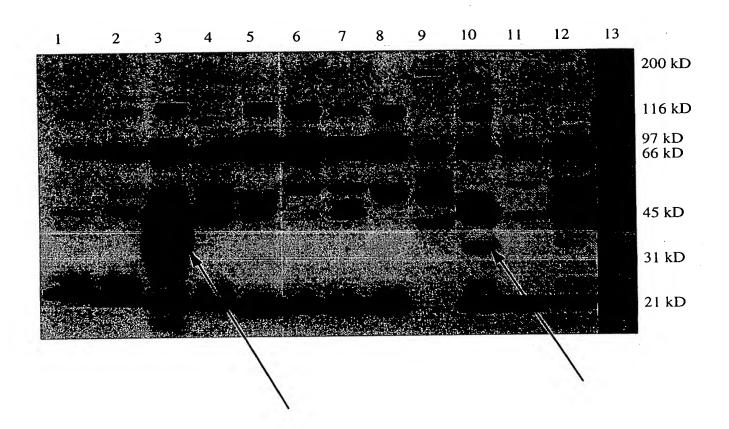
Lane	Tissue	Lane	Tissue
1	Normal Bladder	7	Bladder Cancer
2	Normal Bladder	8	Bladder Cancer
3	Normal Bladder	9	Bladder Cancer
4	Normal Bladder	10	Bladder Cancer
5	Normal Bladder	11	Bladder Cancer
6	Normal Bladder	12	PS116 Bacterial RNA

DODYGOVA.CELEGE

APPROVED O.G. FIG.
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Figure 4



Lane	Tissue	Lane	Tissue
1	Bladder Cancer	8	Breast
2	Bladder	9	Prostate Cancer
3	Bladder Cancer	10	Prostate Cancer
4	Bladder	11	Prostsate Cancer
5	Bladder Cancer	12	ВРН
6	Colon	13	Markers
7	Lung		·

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